5]:	<pre># 1. Generate swiss-roll dataset x,y = datasets.make_swiss_roll(n_samples = 2000) swiss_roll_dataframe = pd.DataFrame(x, columns=['x', 'y', 'z']) swiss_roll_dataframe['output'] = y swiss_roll_dataframe.head(10) x</pre>
;	0 2.136586 3.951359 -10.981817 11.187730 1 11.844955 19.672883 5.330870 12.989270 2 -8.113438 4.256098 -5.942577 10.056943 3 5.467529 17.498402 4.287641 6.948218 4 11.581824 2.103124 6.001622 13.044467 5 -3.873607 5.538540 7.382670 8.337185 6 -2.621947 10.182603 7.748682 8.180261
H b	7 -1.453736 16.491614 -10.763599 10.861326 8 12.573947 14.082619 0.100300 12.574347 9 0.365166 18.595950 14.106561 14.111286 Here we plot out generated swiss roll. This time I made it in color, unfortunately next times I ommited color and I think this is my mistate because I assume it could help me analyze the work better. Probably while writing these markdowns I will add color if I have time. We need to apply a function that reduces amount of dimensions. Since we have 3 dimensions here, I will try n_components = 2 here are atter in the file.
	<pre># 2. apply PCA and plot the data # I assume if we start with 3 columns of training data, we need to make it 2? pca_fit = decomposition.PCA(n_components=2) pca_swiss_roll = pd.DataFrame(pca_fit.fit_transform(swiss_roll_dataframe[['x', 'y', 'z']]), columns=['PCF pca_swiss_roll['output'] = swiss_roll_dataframe['output'] fig = plt.figure(figsize=(13, 13)) ax = fig.add_subplot(111, projection='3d') ax.scatter(x[:, 0], x[:, 1], x[:, 2], c=y, cmap=plt.cm.rainbow) plt.show()</pre>
	15
	5 0 -5 -5
d A A	This is yet another dimension reduction function. Unfortunately there was no single word that I need to plot it, just to find errors. So I didn't plot it, only showed errors. A couple things to note here is 1) constructor of LLE needed 'eigen_solver' parameter and it would fail in a loop without it for some real and 2) I got an unexpected paratemer with a negative error which I assumed to be something interesting. Did I beat machine learning made a 100.000000000000000000000000000000000
m C	Also, from my point of view options of neighbot 1 to 8 were similar at least on the plot, but of course most effective of them is 3 *(or 4 muhahahahahah) Ok, now this is weird. On another run I got more negative errors. I would be please if someone explains this to me. # 3. Apply LLE with 5 neighbours # again we reduce the dimensions into 2 lle = manifold.LocallyLinearEmbedding(n_neighbors=5, n_components=2, n_jobs=-1) lle_swiss_roll = pd.DataFrame(lle.fit_transform(swiss_roll_dataframe[['x', 'y', 'z']]), columns = ['LLE1' lle_swiss_roll['output'] = swiss_roll_dataframe['output'] print(lle.reconstruction error)
	<pre>neighbor_amount = [] errors = [] for nr_of_neighbors in range(2, 16): print(nr_of_neighbors) lle_loc = manifold.LocallyLinearEmbedding(n_neighbors=nr_of_neighbors, n_components=2, n_jobs=-1, eighbor_amount.append(nr_of_neighbors) errors.append(lle_loc.reconstruction_error_) for x, y in zip(neighbor_amount, errors): print(f'neighbor %s with the error of %s' % (x, y))</pre>
	<pre>plt.scatter(neighbor_amount, errors) plt.xlabel('neighbot amount') plt.ylabel('error') plt.show() 2.955875275395633e-17 2 3 4 5 6 7 8</pre>
	9 10 11 12 13 14 15 neighbor 2 with the error of -1.2642328883985104e-14 neighbor 3 with the error of -1.2021613179401072e-14 neighbor 4 with the error of -2.6584849058430206e-15 neighbor 5 with the error of 8.769405302903903e-17 neighbor 6 with the error of 1.9240819886570646e-09 neighbor 7 with the error of 1.6207843957911156e-09
1 1 1 1	neighbor 8 with the error of 3.0702310404986077e-09 neighbor 9 with the error of 3.5747254544005664e-09 neighbor 10 with the error of 8.359382024127963e-09 neighbor 11 with the error of 8.85928312395945e-09 neighbor 12 with the error of 7.015089269694465e-08 neighbor 13 with the error of 8.793319848005186e-08 neighbor 14 with the error of 8.795424019702995e-08 neighbor 15 with the error of 1.0437723769062499e-07
	t is time to apply MDS, and I make my mistake of ignoring the color. Let me write second one with seaborn and fix this.
2]:	<pre># 4. use multi dimensional scaling and visualize the dataset in 2 dimensions mds = manifold.MDS(n_components=2, n_jobs=-1) mds_swiss_roll = pd.DataFrame(mds.fit_transform(swiss_roll_dataframe[['x', 'y', 'z']]), columns = ['MDS1' mds_swiss_roll['output'] = swiss_roll_dataframe['output'] plt.scatter(mds_swiss_roll['MDS1'], mds_swiss_roll['MDS2']) plt.show()</pre>
	15 - 10 - 5 - 0 - -5 - -10 - -15 -
3]:	-15 -10 -5 0 5 10 15 sns.pairplot(data=mds_swiss_roll, hue='output') <seaborn.axisgrid.pairgrid 0x1c99484efa0="" at=""></seaborn.axisgrid.pairgrid>
	Output 6 8 10 10 112 14
	Now, when I have output color on the plot, I can clearly see what is where, it is very interesting if you see the 'output' value together woosition. # 5. apply t-SNE model to the dataset and visualize tsne = manifold.TSNE(n_components=2, n_jobs=-1) tsne_swiss_roll = pd.DataFrame(tsne.fit_transform(swiss_roll_dataframe[['x', 'y', 'z']]), columns = ['TSN tsne_swiss_roll['output'] = swiss_roll_dataframe['output']
6]:	<pre>C:\Users\otaku\AppData\Local\Programs\Python\Python39\lib\site-packages\sklearn\manifold_t_sne.py:780: For arning: The default initialization in TSNE will change from 'random' to 'pca' in 1.2. warnings.warn(C:\Users\otaku\AppData\Local\Programs\Python\Python39\lib\site-packages\sklearn\manifold_t_sne.py:790: For arning: The default learning rate in TSNE will change from 200.0 to 'auto' in 1.2. warnings.warn(plt.scatter(tsne_swiss_roll['TSNE1'], tsne_swiss_roll['TSNE2']) plt.show()</pre>
7]:	-80 -60 -40 -20 0 20 40 Ok, I want to make this in color as well, because this weird figure might seem of a weird shape, but maybe colors will make sense? sns.pairplot(data=tsne_swiss_roll, hue='output') <seaborn.axisgrid.pairgrid 0x1c9947f0e80="" at=""></seaborn.axisgrid.pairgrid>
	Output -75 40 20 12 14
р	Once again colors prove to have more value than the shape. Yes, the figure is not roll shaped as before, but I have clear vision of what point type is closer to what end of the figure. Anyhow, I would like to leave colored seaborn roll as the best one(mds) because it saves shape and point positioning well. So, visually it is my winner. What actual numbers tell us about model comparison is later. # 2.1 import digit dataset with 6 classes
	<pre>digits = datasets.load_digits(n_class=6) print(digits) X, y = digits.data, digits.target n_samples, n_features = X.shape n_neighbors = 30 {'data': array([[0., 0., 5., , 0., 0., 0.],</pre>
3	[0, 0, 1,, 6, 0, 0, 0], [0, 0, 0, 0, 0, 0, 0], [0, 0, 0,, 10, 0, 0, 0], [0, 0, 0,, 10, 0, 0, 0], [0, 0, 0,, 12, 0, 0, 0], [0, 0, 3,, 14, 0, 0], [0, 0, 8,, 16, 0, 0],, [0, 0, 16,, 0, 0, 0], [0, 3, 15,, 11, 5, 0], [0, 0, 0,, 16, 0, 0], [0, 0, 0,, 16, 0, 0], [0, 0, 0,, 16, 0, 0], [0, 0, 0,, 16, 0, 0], [0, 0, 0,, 2, 0, 0], [0, 0, 0,, 2, 0, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2, 0], [0, 0, 0,, 1, 2,, 1, 2, 0], [0, 0, 0, 0,, 1, 2,, 1, 2,, 1, 2], [0, 0, 0, 0,, 1, 1, 1, 0, 0], [0, 0, 0, 0, 0, 0, 0], [0, 0, 0, 0, 0, 0, 0], [0, 0, 0, 0, 0, 0, 0], [0, 0, 0, 0, 0, 0], [0, 0, 0,
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